

## SEQ ID No.1

5'GTTAGCTCAGATTGAACGCTGGCGGCAGGCTTAAACACATGCAAGTCGAGC  
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CGCATACGCCTGAGGGGGAAAGGAGGGGACTCTTCGGAGCCTTCCGCTATT  
AGATGAGCCTGCGTGAGATTAGCTAGTTGGTAGGGTAAAGGCCTACCAAGGC  
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CGGCCAGACTCCTACGGGAGGCAGCAGTGGGGAATATTGGACAATGGGCGC  
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CACTTTCAGGGGTGAGGAAGGGTGATAGGTTAATACGTTATCATCTTGACGTT  
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CTGGCTAGCTAGAGTATGGTAGAGGGGTGTGGAATTCCTGTGTAGCGGTGA  
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CCCTTGTCCTTATTTGCCAGCACGTAATGGTGGGAACTTTAAGGAGACTGCCG  
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AGTAGGGCTACACACGTGCTACAATGGCGTATACAGAGGGCTGCAAGCTAGC  
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CTCGACTCCATGAAGTCGGAATCGCTAGTAATCGTGAATCAGAATGTCACGG  
TGAATACGTTCCCGGCCCTTGACACACCGCCCGTCACACCATGGGAGTTGAT  
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GTCAATGACTGGGGTTGAAGTCTACGCG-3

## SEQ 1D No. 2

1 5'GCCCTTGCTC AGATTGAACG CTGGCGGCAG GCCTAACACA TGCAAGTCGA  
51 GCGGTAGAGA GAAGCTTGCT TCTCTTGAGA GCGGCGGACG GGTGAGTAAT  
101 GCCTAGGAAT CTGCCTGGTA GTGGGGGATA ACGTTCGGAA ACGGACGCTA  
151 ATACCGCATA CGTCCTACGG GAGAAAGCAG GGGACCTTCG GGCCTTGCGC  
201 TATCAGATGA GCCTAGGTCG GATTAGCTAG TTGGTGAGGT AATGGCTCAC  
251 CAAGGCGACG ATCCGTAACT GGTCTGAGAG GATGATCAGT CACACTGGAA  
301 CTGAGACACG GTCCAGACTC CTACGGGAGG CAGCAGTGGG GAATATTGGA  
351 CAATGGGCGA AAGCCTGATC CAGCCATGCC GCGTGTGTGA AGAAGGTCTT  
401 CGGATTGTAA AGCACTTTAA GTTGGGAGGA AGGTTGTAG ATTAATACTC  
451 TGCAATTTTG ACGTTACCGA CAGAATAAGC ACOGGCTAAC TCTGTGCCAG  
501 CAGCCGCGGT AATACAGAGG GTGCAAGCGT TAATCGGAAT TACTGGGCGT  
551 AAAGCGCGCG TAGGTGGTTT GTTAAGTTGG ATGTGAAATC CCCGGGCTCA  
601 ACCTGGGAAC TGCATTCAA ACTGACTGAC TAGAGTATGG TAGAGGGTGG  
651 TGGAATTTCC TGTGTAGCGG TGAAATGCGT AGATATAGGA AGGAACACCA  
701 GTGGCGAAGG CGACCACCTG GACTAATACT GACACTGAGG TGCGAAAGCG  
751 TGGGGAGCAA ACAGGATTAG ATACCTTGT AGTCCACGCC GTAAACGATG  
801 TCAACTAGCC GTTGAAGCC TTGAGCTTTT AGTGGCGCAG CTAACGCATT  
851 AAGTTGACCG CCTGGGGAGT ACGGCCGCAA GGTTAAACT CAAATGAATT  
901 GACGGGGGCC CGCACAAGCG GTGGAGCATG TGGTTTAATT CGAAGCAACG  
951 CGAAGAACCT TACCAGGCCT TGACATCCAA TGAACTTTCT AGAGATAGAT  
1001 TGGTGCCTTC GGGAAACATT AGACAGGTGC TGCATGGCTG TCGTCAGCTC  
1051 GTGTTGTGAA ATGTAAGGGC-3'